IN THE CLAIMS:

Please cancel claim 1 and add new claims 32-60, as follows:

Claims 1-31 (Canceled).

- 32. (New) A chimeric nonhuman/human antibody or a fragment, a region, or a derivative thereof, wherein the chimeric antibody:
 - (a) binds to lipoteichoic acid of Gram positive bacteria;
 - (b) enhances opsonization of Gram positive bacteria by 75% or more over background; and
 - (c) confers a statistically significant enhancement of survival in a lethal animal model.
- 33. (New) A chimeric nonhuman/human antibody or a fragment, a region, or a derivative thereof, wherein the chimeric antibody:
 - (a) binds to lipoteichoic acid of Gram positive bacteria; and
 - (b) enhances the opsonization of Gram positive bacteria by 75% or more over background.
- 34. (New) The antibody of claim 33, wherein the chimeric antibody binds to lipoteichoic acid at a level that is twice the background or greater.
- 35. (New) The antibody of claim 33, wherein the chimeric antibody further recognizes a peptide sequence chosen from:

WRMYFSHRHAHLRSP(SEQ ID NO 1); and WHWRHRIPLQLAAGR(SEQ ID NO 2).

36. (New) The antibody of claim 35, wherein the chimeric antibody binds to lipoteichoic acid at a level that is twice the background or greater.

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- 37. (New) The antibody of claim 33, wherein the chimeric antibody is an IgG isotype or an IgM isotype.
- 38. (New) The antibody of claim 33, wherein the fragment of the chimeric antibody is chosen from at least one of Fab, Fab', F(ab')₂, and SFv.
- 39. (New) The antibody of claim 33, wherein the chimeric antibody enhances survival in a lethal animal model by 10% or more over animals not treated with the antibody.
- 40. (New) The antibody of claim 39, wherein the chimeric antibody further recognizes a peptide sequence chosen from:

WRMYFSHRHAHLRSP(SEQ ID NO: 1); and WHWRHRIPLQLAAGR(SEQ ID NO: 2).

- 41. (New) The antibody of claim 33, wherein the chimeric antibody's Complementarity Determining Regions correspond to at least one of the Complementarity Determining Regions of Figure 12.
- 42. (New) The antibody of claim 41, wherein the Complementarity Determining Region amino acid sequences are at least 70% homologous to the Complementarity Determining Region amino acid sequences chosen from at least one of:
 - (a) amino acids 31-35 of SEQ ID NO. 87, amino acids 50-68 of SEQ ID NO.

87, and amino acids 101-112 of SEQ ID NO. 87; and wherein the Complementarity Determining Region amino acid sequences are at least 70% homologous to the Complementarity Determining Region amino acid sequences further chosen from at least one of:

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- (b) amino acids 24-33 of SEQ ID NO. 89, amino acids 49-55 of SEQ ID NO. 89, and amino acids 88-96 of SEQ ID NO. 89.
- 43. (New) A chimeric nonhuman/human antibody or a fragment, a region, or a derivative thereof, comprising at least part of a human immunoglobulin constant region and at least part of a nonhuman immunoglobulin variable region having specificity to lipoteichoic acid of Gram positive bacteria, wherein the chimeric antibody:
 - (a) binds to lipoteichoic acid of Gram positive bacteria; and
 - (b) enhances the opsonization of Gram positive bacteria by 75% or more over background.
- 44. (New) The antibody of claim 43, wherein the chimeric antibody binds to lipoteichoic acid at a level that is twice the background or greater.
- 45. (New) The antibody of claim 43, wherein the chimeric antibody further recognizes a peptide sequence chosen from:

WRMYFSHRHAHLRSP (SEQ ID NO 1); and WHWRHRIPLQLAAGR (SEQ ID NO 2).

- 46. (New) The antibody of claims 43, 44, or 45, wherein the nonhuman immunoglobulin variable region is the nonhuman immunoglobulin variable region of monoclonal antibody 96-110.
- 47. (New) The antibody of claim 43, wherein the human immunoglobulin constant region is subclass IgG1.
- 48. (New) The antibody of claim 43, wherein the chimeric antibody is an IgG isotype or an IgM isotype.

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- 49. (New) The antibody of claim 43, wherein the fragment of the chimeric antibody is chosen from at least one of Fab, Fab', F(ab')₂, and SFv.
- 50. (New) The antibody of claim 43, wherein the chimeric antibody enhances survival in a lethal animal model by 10% or more over animals not treated with the antibody.
- 51. (New) The antibody of claim 50, wherein the chimeric antibody further recognizes a peptide sequence chosen from:

WRMYFSHRHAHLRSP(SEQ ID NO: 1); and WHWRHRIPLQLAAGR(SEQ ID NO: 2).

- 52. (New) The antibody of claim 43, wherein the chimeric antibody's Complementarity Determining Regions correspond to at least one of the Complementarity Determining Regions of Figure 12.
- 53. (New) The antibody of claim 52, wherein the Complementarity Determining Region amino acid sequences are at least 70% homologous to the Complementarity Determining Region amino acid sequences chosen from at least one of:
 - (a) amino acids 31-35 of SEQ ID NO. 87, amino acids 50-68 of SEQ ID NO.

87, and amino acids 101-112 of SEQ ID NO. 87; and wherein the Complementarity Determining Region amino acid sequences are at least 70% homologous to the Complementarity Determining Region amino acid sequences further chosen from at least one of:

(b) amino acids 24-33 of SEQ ID NO. 89, amino acids 49-55 of SEQ ID NO. 89, and amino acids 88-96 of SEQ ID NO. 89.

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- 54. (New) A composition comprising at least one of the antibodies of claims 32, 33, or 43 and a pharmaceutically acceptable carrier.
- 55. (New) A composition comprising a fragment, a region, or a derivative of at least one of the antibodies of claims 32, 33, or 43, and a pharmaceutically acceptable carrier.
- 56. (New) The composition of claim 55, wherein the fragment of the antibody is chosen from at least one of Fab, Fab', F(ab')₂, and SFv.
- 57. (New) The composition of 55, wherein the region is a Complementarity Determining Region.
- 58. (New) The composition of 55, wherein the derivative is comprised of proteins or peptides encoded by truncated or modified antibody genes.
- 59. (New) A murine monoclonal antibody, wherein the monoclonal antibody comprises monoclonal antibody 96-110.
- 60. (New) The antibody of claim 59, wherein a fragment of the monoclonal antibody is chosen from at least one of Fab, Fab', F(ab')₂, and SFv.

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